

INFORMATION DISCLOSURE STATEMENT BY APPLICANT <i>(Use as many sheets as necessary)</i>				Complete if Known	
				Application Number	10/606,618
				Filing Date	June 26, 2003
				First Named Inventor	Judd, et al.
				Art Unit	1645
				Examiner Name	S. Devi
Sheet	1	of	2	Attorney Docket Number	UM/SBC147BUSA

U.S. PATENT DOCUMENTS					
Examiner Initials*	Cite No. ¹	Document Number	Publication Date MM-DD-YYYY	Name of Patentee or Applicant of Cited Document	Pages, Columns, Lines, Where Relevant Passages or Relevant Figures Appear
		Number-Kind Code ² (if known)			
	AC	US-6,348,185	02-19-2002	Washington University School of Medicine	
	AD	US-2005-0074458	04-07-2005	Judd, et al.	
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FOREIGN PATENT DOCUMENTS						
Examiner Initials*	Cite No. ¹	Foreign Patent Document	Publication Date MM-DD-YYYY	Name of Patentee or Applicant of Cited Document	Pages, Columns, Lines, Where Relevant Passages or Relevant Figures Appear	T ⁶
		Country Code ³ - Number ⁴ -Kind Code ⁵ (if known)				
	AE	EP – 1 535 928	07-16-2008	The University of Montana		
	AF	WO – 94/12641	06-09-1994	Connaught Laboratories Limited		

Examiner Signature	Date Considered
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NON PATENT LITERATURE DOCUMENTS				
Examiner Initials*	Cite No. ¹	Include the name of the author (in CAPITAL LETTERS), title of the article (when appropriate), title of the item, (book, magazine, journal, serial, symposium, catalog, etc.), date, page(s), volume-issue number(s), publisher, city and/or country where published	T ²	
	AG	BHASIN, et al. Infect. Immun., Vol. 69, pp. 5031-5036, (August 2001)		
	AH	HARLOW et al. , In: Antibodies: A Laboratory Manual, Cold Spring Harbor Laboratory, Chapter 5, pg. 76, (1988)		
	AI	KAWARABAYASHI et al. , Complete Sequence and Gene Organization of the Genome of a Hyper-Thermophilic Archaeobacterium, <i>Pyrococcus Horikoshii</i> OT3DNA Res. 5: 55-76, (1998)		
	AJ	Protein Sequence on STN, pg. 12, Sequence alignment provided by Examiner in Office Action dated August 5, 2005 for US Patent Application No. 10/606,618 from STN International		
	AK	SKOLNICK et al. , From genes to protein structure and function: novel applications of computational approaches in the genomic era, Trends in Biotechnology, 18(1):34-39, (January, 2000)		

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